## **Amendments to the Sequence Listing**

Please replace the originally filed sequence listing with the amended sequence listing attached to this amendment. All amendments relate to improper numbering or improper naming of the modifications to the sequences. Thus, no new matter is added through any of these amendments.

The following amendments were made to the sequence listing:

			_		
Seq	Originally read		As amended reads		Reason for
No.					amendment;
					basis in
					specification
26	<220>		<220>		misnumbered;
	I	MOD_RES	l .	MOD_RES	see Table 1,
	<222>	$(5) \dots (5)$		$(8) \dots (8)$	Compound 26
		AMIDATION		AMIDATION	
65	<220>		<220>		end of phrase ("D
		MOD_RES	<221>	MOD_RES	form") omitted;
	<222>	$(1)\dots(1)$	<222>	(1)(1)	divided into two
	<223>	N-(2-	<223>	N-(2-	entries;
		napthylenesulfonyla		napthylenesulfonyla	see Table 1,
		mino-4-oxo-butyryl)		mino-4-oxo-	Compound 65
		substituted, D		butyryl) substituted	
65			<220>		end of phrase ("D
			<221>	MOD_RES	form") omitted;
			<222>	(1)(1)	divided into two
			<223>	D form	entries;
					see Table 1,
					Compound 65
78	<220>		<220>		omitted identity
	<221>	MOD_RES	<221>	MISC_FEATURE	of Xaa residue
		(9)(9)	<222>	(9)(9)	that was reduced;
	<223>	Reduced from amino	<223>		see Table 1,
		acid to amino		from amino acid to	Compound 78
		alcohol		amino alcohol	*
94		·	<220>		omitted disulfide
1				DISULFID	description;
				(3)(9)	see Table 1,
				., .,	Compound 94
116	<220>		<220>		omitted identity
	i —— - :	MOD_RES		MISC_FEATURE	of Xaa residue
1	<222>			(10)(10)	that was reduced;
[	<223>		<223>	Xaa = Ser reduced	see Table 1,
		acid to amino		from amino acid to	Compound 116
		alcohol		amino alcohol	
	<u> </u>		L		

Seq No.	Originally read		As amended reads		Reason for amendment; basis in specification
120	<220>	MOD DEG	<220>	MICC FEATURE	omitted identity
		MOD_RES		MISC_FEATURE	of Xaa residue
		(11)(11)		(11)(11)	that was reduced;
	<223>	reduced from amino	<223>	Xaa = Pro reduced from amino acid to	see Table 1,
		acid to amino			Compound 120
122	<220>	alcohol	<220>	amino alcohol	omitted identity
122		MOD DEC		MISC_FEATURE	of Xaa residue
		MOD_RES (11)(11)		(11)(11)	that was reduced;
		reduced from amino		Xaa = Pro reduced	see Table 1,
	\2237	acid to amino	\2237	from amino acid to	Compound 122
		alcohol		amino alcohol	Compound 122
159	<220>	uiconor			incorrectly listed
107		MOD_RES			in Table 1 as
		$(7)$ $\dots$ $(7)$			amidated; should
		AMIDATION			be acid form
166	<220>		<220>		misnumbered;
	<221>	DISULFID	<221>	DISULFID	see Table 1,
	<222>	(3)(8)	<222>	(3) (9)	Compound 166
166	<220>		<220>		misnumbered;
		MOD_RES		MOD_RES	see Table 1,
	1	(5)(5)		(6)(6)	Compound 166
	<del></del>	D form	<del></del>	D form	
166	<220>		<220>		misnumbered;
	1	MOD_RES	1	MOD_RES	see Table 1,
	<222>	(8)(8)	<222>	(9)(9)	Compound 166
100	<223>	AMIDATION	<223>	AMIDATION	
182	<220>	MOD DEC	<220>	MOD DEC	incorrectly listed
	L.	MOD_RES	1	MOD_RES	as 4-fluoro rather
		(3)(3)	1	(3)(3)	than 4-chloro;
	<223>	· · · · · · · · · · · · · · · · · · ·	<223>	•	see Table 1,
		D form	<u> </u>	D form	Compound 182